

# SEQUENCE LISTING

<110> Napier, Johnathan A.

<120> Polyunsaturated Fatty Acid (PUFA) Elongase from *Caenorhabditis elegans*

<130> 76/7

<140> PCT/GB00/01035

<141> 2000-03-20

<160> 22

<170> PatentIn Ver. 2.1

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 Phe Ile Ile Leu Arg Lys Lys Pro Leu Met Phe Leu His Trp Tyr His  
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 His Val Leu Thr Met Asn Tyr Ala Phe Met Ser Phe Glu Ala Asn Leu  
       145                          150                          155                          160  
 Gly Phe Asn Thr Trp Ile Thr Trp Met Asn Phe Ser Val His Ser Ile  
                           165                          170                          175  
 Met Tyr Gly Tyr Tyr Met Leu Arg Ser Phe Gly Val Lys Val Pro Ala  
           180                          185                          190  
 Trp Ile Ala Lys Asn Ile Thr Thr Met Gln Ile Leu Gln Phe Val Ile  
           195                          200                          205  
 Thr His Phe Ile Leu Phe His Val Gly Tyr Leu Ala Val Thr Gly Gln  
           210                          215                          220  
 Ser Val Asp Ser Thr Pro Gly Tyr Tyr Trp Phe Cys Leu Leu Met Glu  
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 Ile Ser Tyr Val Val Leu Phe Gly Asn Phe Tyr Tyr Gln Ser Tyr Ile  
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**FBI**

12

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 Asp Ala Glu Gly Arg Lys Phe Phe Ala Asp His Phe Asp Val Thr Ile  
 35 40 45  
 Gln Ala Ser Ile Leu Tyr Met Val Val Val Phe Gly Thr Lys Trp Phe  
 50 55 60  
 Met Arg Asn Arg Gln Pro Phe Gln Leu Thr Ile Pro Leu Asn Ile Trp  
 65 70 75 80  
 Asn Phe Ile Leu Ala Ala Phe Ser Ile Ala Gly Ala Val Lys Met Thr  
 85 90 95  
 Pro Glu Phe Phe Gly Thr Ile Ala Asn Lys Gly Ile Val Ala Ser Tyr  
 100 105 110  
 Cys Lys Val Phe Asp Phe Thr Lys Gly Glu Asn Gly Tyr Trp Val Trp  
 115 120 125  
 Leu Phe Met Ala Ser Lys Leu Phe Glu Leu Val Asp Thr Ile Phe Leu  
 130 135 140  
 Val Leu Arg Lys Arg Pro Leu Met Phe Leu His Trp Tyr His His Ile  
 145 150 155 160  
 Leu Thr Met Ile Tyr Ala Trp Tyr Ser His Pro Leu Thr Pro Gly Phe  
 165 170 175  
 Asn Arg Tyr Gly Ile Tyr Leu Asn Phe Val Val His Ala Phe Met Tyr  
 180 185 190  
 Ser Tyr Tyr Phe Leu Arg Ser Met Lys Ile Arg Val Pro Gly Phe Ile  
 195 200 205  
 Ala Gln Ala Ile Thr Ser Leu Gln Ile Val Gln Phe Ile Ile Ser Cys  
 210 215 220  
 Ala Val Leu Ala His Leu Gly Tyr Leu Met His Phe Thr Asn Ala Asn  
 225 230 235 240  
 Cys Asp Phe Glu Pro Ser Val Phe Lys Leu Ala Val Phe Met Asp Thr  
 245 250 255



